

SEQUENCE LISTING

<110> Riechmann, Lutz

Kristensen, Peter

Jestin \ Jean-Luc

Winter, Gregory

<120> Selection System

<130> 8039/1090

<140> 09/710,444

<141> 2000-11-10

<150> GB 9810223.9

<151> 1998-05-13

<150> GB 9810228.8

<151> 1998-05-13

<150> PCT/GB99/01526

<151> 1999-05-13

<160> 78

<170> PatentIn version 3.1

<210> 1

```
<211> 17
```

<220>

<223> Synthetic linker peptide sequence with protease recognition sites

<400> 1

Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu Gly Arg Gly Ala His

012

5

10

15

Glu

<211> 57

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(57)

	<400>	2	
	ggcacc	ctca gaacggtacc ccaccctcag aggccggctg ggccgccacc ctcagag	57
	<210>	3	
	<211>	89	
	<212>	DNA	
	<213>	Artificial	
)			
_	<220>		
	<221>	misc_feature	
	<222>	(1)(89)	
	<223>	Synthetic PCR primer for vector construction/screening	
	<400>	3	
	ggtggc	ggcc cagceggcct ttetgagggg tegaetatag aaggaegagg geecagegaa	60
	ggaggt	gggg tacccccttc tgagggtgg	89
	<210>	4	
	<211>	89	
	<212>	DNA	
	<213>	Artificial	

<223> Synthetic PCR primer for vector construction/screening

<220>		
<221>	misc_feature	
<222>	(1)(89)	
<223>	Synthetic PCR primer for vector construction/screening	
<400>	4	
ccaccc	tcag aagggggtac cccacctcct tcgctgggcc ctcgtccttc tatagtcgac	60
ccctca	gaaa ggccggctgg gccgccacc	89

<222> (1)..(24)

<223> Synthetic PCR primer for vector construction/screening

<400> 5
gcgatggttg ttgtcattgt cggc

```
<210> 6
```

- <220>
- <221> misc_feature
- <222> (1)..(24)
- <223> Synthetic PCR primer for vector construction/screening
- <400> 6

aaaagaaacg caaagacacc acgg

24

<210>

- <212> DNA
- <213> Artificial
- <220>
- <221> misc_feature
- <222> (1)..(23)
- <223> Synthetic PCR primer for vector construction/screening
- <400> 7

cctcctgagt acggtgatac acc

```
<210> 8
```

<220>

<223> Synthetic PCR primer used to screen for recombinant clones

<400> 8

gtaaattcag agactgcgct ttcc

24

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(26)

<223> Synthetic PCR primer used to screen for recombinant clones

	<400>	9	
	attttc	ggtc atagccccct tattag	26
	<210>	10	
	<211>	65	
	<212>	DNA	
	<213>	Artificial	
	<220>		
	<221>	misc_feature	
	<222>	(1)(65)	
	<223>	Synthetic PCR primer recognizing FLAG tag nucleotide sequence	
		·	
	<400>	10	
,	caaacg	ggcg gccgcagact acaaggatga cgacgacaag gaaactgttg aaagttgttt	60
	agcaa		65

<210> 11

<211> 51

<212> DNA

<213> Artificial

<220>

<221> misc_feature

- <222> (1)..(51)
- <223> Synthetic PCT primer used to change codon usage in recombinant cl ones
- <400> 11

cccctcagaa aggccggctg ggccgccgcc agcattgaca ggaggttcag g 51

- <210> 12
- <211> 52
- <212> DNA
- <213> Artificial
- <220>
- <221> misc_feature
- <222> (1)..(52)
- <223> Synthetic PCT primer used to change codon usage in recombinant cl ones
- <400> 12

gaaggaggtg gggtacccgg ttccgagggt ggttccggtt ccggtgattt tg 52

- <210> 13
- <211> 36
- <212> DNA

	<213>	Artificial	
	222		
	<220>		
	<221>	misc_feature	
	<222>	(1)(36)	
	<223>	Synthetic PCR primer for vector construction/screening	
	<400>	13	
	ccctcg	gaac cggtacccca gctgcttcgt gggccc	36
	<210>	14	
7	<211>		
	<212>	DNA	
	<213>	Bacillus amyloliquefaciens	
	<400>	14	
	ctggcg	gcgg cccagccggc cctgcacagg ttatcaacac gtttgac	47
	<210>	15	
	<211>	43	
	<212>	DNA	
	<213>	Bacillus amyloliquefaciens	

43

ctcggaaccg gtacctctga tttttgtaaa ggtctgataa gcg

<400> 15

<210> 16

<211> 44

<212> DNA

<213> Gallus gallus

<400> 16

ggcggcccag ccggcctttc tctctctgac gaggacttca aggc

44

<210> 17

<211> 41

<212> DNA

]<213> Gallus gallus

<400> 17

cctcggaacc ggtaccgaag agtcctttct ccttcttgag g

41

<210> 18

<211> 18

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(18)

	<400>	18	
	tacgcc	aagc ttgcatgc	18
	<210>	19	
	<211>	17	
	<212>	DNA	
	<213>	Artificial	
	<220>		
		misc_feature	
) _		(1)(17)	
	<223>	Synthetic PCR primer used for library construction	
	<400>	19	
		ctgg gccatgg	17
	<210>	20	
	<211>	17	
	<212>	DNA	

<223> Synthetic PCR primer used for library construction

<213> Artificial

<220>

```
<221> misc_feature
<222> (1)..(17)
<223> Synthetic PCR primer used for library construction
<400> 20
gattacgcca agctttg
                                                                     17
<210> 21
<211> 126
<212> DNA
<213> Erwinia chrysanthemi
<220>
      misc_feature
<223> n at positions 23, 24, 29, 55, 56, 81, 97, 101, and 102 can be G,
       A, T or C
<220>
<221> misc_feature
<222> (23)..(23)
<223> n at position 23 can be G, A, T or C
<220>
```

<221> misc_feature

```
<222> (24)..(24)
<223> n at position 24 can be G, A, T or C
<220>
<221> misc_feature
<222> (29)..(29)
<223> n at position 29 can be G, A, T or C
<220>
<221> misc_feature
<222> (55)..(55)
<223> n at position 55 can be G, A, T or C
<220>
<221> misc_feature
<222> (56)..(56)
<223> n at position 56 can be G, A, T or C
<220>
<221> misc_feature
```

<223> n at position 81 can be G, A, T or C

<222> (81)..(81)

```
<220>
<221> misc_feature
<222> (97)..(97)
<223> n at position 97 can be G, A, T or C
<220>
<221> misc feature
```

<222> (101)..(101)

<223> n at position 101 can be G, A, T or C

<220>
<221> misc_feature

(102)..(102)

<223> n at position 102 can be G, A, T or C

<400> 21
gattacgcca agcttgcatg cannddctnt dtcaaggaga cagtcataat garrnnbcta 60
ttgsyaayrs yasyasyagb nttgttatta ctcsyanycv nncygdccat ggcccaggtg 120

<210> 22 <211> 117

cagctg

```
<212> DNA
<213> Bacteriophage M13mp18
<220>
<221> misc_feature
<222> (18)..(18)
<223> Nucleotide at position 18 can be G, A, T or C.
<220>
<221> misc_feature
<222> (19)..(19)
<223> Nucleotide at position 19 can be G, A, T or C.
<220>
<221> misc_feature
<222> (20)..(20)
<223> Nucleotide at position 20 can be G, A, T or C.
<220>
<221> misc_feature
<222> (21)..(21)
```

<223> Nucleotide at position 21 can be G, A, T or C.

<400> 22

	gattac	gcca agctttgnnn ncttttttww ggagattttc aacrtgaraa rattattatt	60
	csyaat	tsyt ttagttsyts ytttetwtgy ggyccageeg gecatggeec aggtgea 1	L 1 7
	<210>	23	
	<211>	18	
	<212>	DNA	
	<213>	Artificial sequence	
	<220>		
	<223>	Synthetic PCR primer for vector construction	
	<400>	23	
	ctttat	gett ceggeteg	18
•	2		
4			
	<210>	24	
	<211>	17	
	<212>	DNA	

<213> Artificial

<221> misc_feature

<222> (1)..(17)

<220>

<223> Synthetic PCT primer for library construction

```
<400> 24
```

cggccccatt cagatcc

17

- <210> 25
- <211> 50
- <212> DNA
- <213> Artificial
- <220>
- <221> misc_feature
- <222> (1)..(50)
- <223> Randomized E. chrysanthemi pelB sequence

112

<400> 25

aagcttgcat gcaaattcta tdtcaaggag acagttataa tgaaatacct

- <210> 26
- <211> 50
- <212> DNA
- <213> Artificial
- <220>
- <221> misc_feature
- <222> (1)..(50)
- <223> Randomized E. chrysanthemi pelB sequence

```
<220>
<221> misc_feature
<222> (14)..(14)
<223> n at position 14 can be G, A, T or C.

<220>
<221> misc_feature
<222> (15)..(15)
<223> n at position 15 can be G, A, T or C.
```

<220>
<221> misc_feature
<222> (45)..(45)
<223> n at position 45 can be G, A, T or C.

<220>
<221> misc_feature

```
<222> (46)..(46)
```

<223> n at position 46 can be G, A, T or C.

<400> 26

aagcttgcat gcannddctn tdtcaaggag acagtcataa tgarrnnbct

50

<210> 27

<211> 50

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

<400> 27

aagcttgcat gcagcatctc tdgcaaggag acagtcataa tgaagacgct

50

<210> 28

<211> 50

<212> DNA

<213> Artificial

```
<220>
```

<221>	misc_	_feature

<223> Randomized E. chrysanthemi pelB sequence

<400> 28

aagcttgcat gcacgggctg tdtcaaggag acagtcataa tgagagggct

50

<210> 29

<211> 50

<212> DNA

<213> Artificial



misc_feature

<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

<400> 29

aagcttgcat gcaccagctc tdtcaaggag acagtcataa tgaggcggct

50

<210> 30

<211> 55

<212> DNA

```
<213> Artificial

<220>

<221> misc_feature

<222> (1)..(55)
```

<223> Randomized E. chrysanthemi pelB sequence

<400> 30

attectaacg geageegetg gattgttatt actegeggee eageeggeea tggee

55

```
<210> 31
<211> 55
<212> DNA
<213> artificial
```

<220>

<221> misc_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence

<220>

<221> misc_feature

<222> (22)..(22)

<223> n at position 22 can be G, A, T or C.

```
<220>
<221> misc_feature
<222> (38)..(38)
<223> n at position 38 can be G, A, T or C.

<220>
<221> misc_feature
```

<220>

<221> misc_feature

/222> (43)..(43)

<222> (42)..(42)

<223> n at position 43 can be G, A, T or C.

<223> n at position 42 can be G, A, T or C.

<400> 31

attgsyaayr syasyasyag bnttgttatt actcsyanyc vnncygdcca tggcc

55

<210> 32

<211> 55

<212> DNA

<213> Artificial

```
<220>
<221> misc_feature
<222> (1)..(55)
<223> Randomized E. chrysanthemi pelB sequence
<400> 32
attgcyaatg gtactgtyag gattgttatt actcccaccc ggtccgtcca tggcc
<210> 33
<211> 55
<212> DNA
<213> artificial
<220>
      misc_feature
<222> (1)..(55)
```

55

<223> Randomized E. chrysanthemi pelB sequence

<400> 33

attgcyaatg ctagtgcyag ggttgttatt actcccaatc gcgccggcca tggcc 55

<210> 34

<211> 54

<212> DNA

```
<213> Artificial
```

<223> Randomized E. chrysanthemi pelB sequence

<220>

<223> n at position 22 can be G, A, T or C.

<220>

<221> misc_feature

<222> (43)..(43)

<223> n at position 43 can be G, A, T or C.

<220>

<221> misc_feature

<222> (44)..(44)

<223> n at position 44 can be G, A, T or C.

<400> 34

attggtaata gcagcagtag bnttgttagg actcgcaccc ccnncyadcc atgg

- <210> 35
- <211> 22
- <212> PRT
- <213> Erwinia chrysanthemi
- <400> 35

Met Lys Tyr Leu Leu Pro Thr Ala Ala Gly Leu Leu Leu Leu Ala

1 5 10 15

Ala Gln Pro Ala Met Ala

() 12

- <210> 36
- <211> 22
- <212> PRT
- <213> Artificial
- <220>
- <221> MISC_FEATURE
- <222> (1)..(22)
- <223> Randomized E. chrysanthemi pelB sequence

<400> 36

Met Lys Thr Leu Ala Met Val Leu Val Gly Gly Pro Pro Gly Pro Ser

1 5 10

15

Ala Gln Pro Ala Met Ala

20

<210> 37

<211> 21

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(21)

<223> Randomized E. chrysanthemi pelB sequence

<400> 37

Met Arg Gly Leu Ala Met Leu Val Ala Gly Gly Pro Ile Ala Pro Ala

1 5 10 15

Gln Pro Ala Met Ala

20

```
<210> 38
```

<211> 23

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(23)

<223> Randomized E. chrysanthemi pelB sequence

A (2 < 400 > 38

Met Arg Arg Leu Val Pro Ile Thr Ala Ala Val Gly Leu Leu Ala Pro

1 5 10 15

Pro Thr Gln Pro Ala Met Ala

20

<210> 39

<211> 50

<212> DNA

<213> Artificial

```
<220>
<221> misc_feature
<222> (1)..(50)
<223> Randomized bacteriophage M13 g3 sequence
<400> 39
aagctttgga cgcttttttt tggagatttt caacgtgaaa aaattattat
<210> 40
<211> 50
<212>
<220>
<221> misc_feature
```

<220>
<221> misc_feature
<222> (1)..(50)
<223> Randomized bacteriophage M13 g3 sequence

<223> n at position 9 is can be G, A, t or C.

<222> (9)..(9)

```
<220>
<221> misc_feature
<222> (10)..(10)
<223> n at position 10 is can be G, A, t or C.

<220>
<221> misc_feature
<222> (11)..(11)
<223> n at position 11 is can be G, A, t or C.

<220>
<221> misc_feature
<222> (11)..(11)
<223> n at position 11 is can be G, A, t or C.
```

<400> 40

aagctttgnn nncttttttw wggagatttt caacrtgara arattattat

50

<210> 41

<211> 50

<212> DNA

<213> Artificial

<220>

```
<221> misc_feature
```

<223> Randomized bacteriophage M13 g3 sequence.

<400> 41

aagctttggg gccttttttt aggagatttt caacatgaga agattattat

50

<211> 50

<212> DNA

<213> Artificial

<220>

≪221> misc_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

<400> 42

tcgcaattcc tttagttgtt cctttctatg cggcccagcc ggccatggcc

50

<210> 43

<211> 50

<212> DNA

<213> Artificial

	<220>		
	<221>	misc_feature	
	<222>	(1)(50)	
	<223>	Randomized bacteriophage M13 g3 sequence	
	<400>	43	
	tcsyaa	ttsy tttagttsyt sytttctwtg yggyccagcc ggccatggcc	50
	<210>	44	
	<211>	50	
	<212>	DNA	
)	چ ²¹³ >	Artificial	
	<220>		
	<221>	misc_feature	
	<222>	(1)(50)	
	<223>	Randomized bacteriophage M13 g3 sequence	
	<400>	44	
	tcctaa	ttcc tttagttgtt gctttctatg tggtccagcc ggccatggcc	50
	<210>	45	

<211> 22

- <212> PRT
- <213> Artificial
- <220>
- <221> MISC_FEATURE
- <222> (1)..(22)
- <223> Randomized bacteriophage M13 g3 sequence
- <400> 45

Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ala

1 5 10 15

Ala Gln Pro Ala Met Ala

- <210> 46
- <211> 22
- <212> PRT
- <213> Artificial
- <220>
- <221> MISC_FEATURE
- <222> (1)..(22)
- <223> Randomized bacteriophage M13 g3 sequence

<400> 46

Met Arg Arg Leu Leu Leu Ala Pro Pro Val Ala Val Pro Phe Tyr Val

1 5 10 15

Val Gln Pro Ala Met Ala

20

<210> 47

<211> 18

<212> DNA

<213> artificial

<220>

<221> misc_feature

<222> (1)..(18)

<223> Synthetic oligonucleotide primer used as substrate for Stoffel fr agment of Thermus aquaticus DNA polymerase I

<400> 47

tttcgcaaga tgtggcgt 18

```
<210> 48
```

<220>

<223> Synthetic primer used as substrate for Stoffel fragment of Thermu s aquaticus DNA polymerase I

<400> 48

gcgaagatgt gg

12

<210> 49

<211> 30

<212> DNA

<213> artificial

<220>

<221> misc_feature

<222> (1)..(30)

<223> Synthetic oligonucleotide primer used as substrate for Thermus aquaticus DNA polymerase I

```
<400> 49
```

aaatacaaca ataaaacgcc acatcttgcg

30

- <210> 50
- <211> 20
- <212> DNA
- <213> Artificial
- <220>
- <221> misc_feature
- <222> (1)..(20)
- <223> Synthetic oligonucleotide sequence insert containing PstI restric tion site and frame shift for H102A mutant barnase fusion construct fused to p3 gene of phage fd-3.

A12

<400> 50

ctgcaggcgg tgcggccgca

- <210> 51
- <211> 24
- <212> DNA
- <213> artificial
- <220>
- <221> misc_feature

```
<222> (1)..(24)
<223> Synthetic oligonucleotide used for random priming

<220>
<221> misc_feature
<222> (19)..(19)
<223> n at position 19 can be G, A, T or C.

<220>
<221> misc_feature
<222> (20)..(20)
<223> n at position 20 can be G, A, T or C.
```

<220>
<221> misc_feature
<222> (22)..(22)
<223> n at position 22 can be G, A, T or C.

```
<220>
<221> misc_feature
<222> (23)..(23)
<223> n at position 23 can be G, A, T or C.
```

gagcctgcag agctcaggnn nnnn

24

Q12

<210> 52 <211> 23

<212> DNA

<213> artificial

<220>

<221> misc_feature

<222> (1)..(23)

<223> Synthetic PCR primer used to re-amplify randomly amplified E. col i genomic DNA sequences.

cgtgcgagcc tgcagagctc agg

23

<210> 53

<211> 45

<212> PRT

<213> artificial

<220>

<221> MISC_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

UZ<400> 53

Leu Gln Ser Ser Gly Asp Cys Val Ile Ser Asp Thr Cys Ile Ala Gly

1

5

10

15

Met Ala Glu Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val

20

25

30

Gly Leu Thr Ile Thr Val Thr Pro Cys Leu Ser Ser Ala

35

40

- <210> 54
- <211> 44
- <212> PRT
- <213> artificial
- <220>
- <221> MISC_FEATURE
- <222> (1)..(44)
- <223> Barstar binding barnase-p3 fusion insert

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1

5

10

15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

20

25

30

Pro Ser Ser Ala Thr Ile His Cys Leu Ser Ser Ala

35

40

<210> 55

<211> 40

- <212> PRT
- <213> artificial
- <220>
- <221> MISC_FEATURE
- <222> (1)..(40)
- <223> Barstar binding barnase-p3 fusion insert
- <400> 55

Leu Gln Ser Ser Gly Asp Ser Ala Gly Cys Lys Asn Met Thr Gly Gly

1 5 10 15

Q12

Arg Leu Tyr Ala His Thr Leu Glu Ala Ile Ile Pro Gly Phe Ala Val

20 25 30

Ser Ala Pro Ala Cys Glu Pro Ala

- 35 40
- <210> 56
- <211> 33
- <212> PRT
- <213> artificial

```
<220>
```

- <221> MISC_FEATURE
- <222> (1)..(33)
- <223> Barstar binding barnase-p3 fusion insert

1

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

10

15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

20

5

25

30

(12

Ala

<210> 57

<211> 44

<212> PRT

<213> artificial

<220>

<221> MISC_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 57

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

20 25 30

Pro Ser Ser Ala Thr Val Gln Cys Leu Ser Ser Ala

35 40

<210> 58

<211> 41

<212> PRT

<213> artificial

<220>

<221> MISC_FEATURE

<222> (1)..(41)

<223> Barstar binding barnase-p3 fusion insert

Leu Gln Ser Ser Gly Lys Ile Val Gln Ala Gly Ala Asn Ile Gln Asp

1 5 10 15

Gly Cys Ile Met His Gly Tyr Cys Asp Thr Asp Thr Ile Val Gly Glu
20 25 30

Asn Gly His Ile Gly Leu Ser Ser Ala

35 40

<210> 59

<211> 45

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

<400> 59

Leu Gln Ser Ser Gly Val Cys Val Ile Ser Asp Thr Cys Ile Ala Gly

1 5 10 15

Thr Ala Glu Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val $\dot{}$

20 25 30

Gly His Thr Ile Thr Glu Thr Pro Cys Leu Ser Ser Ala

35 40 45

<210> 60

<211> 44

<212> PRT

<213> artificial

<220>

<221> MISC_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 60

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

20

25

30

Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala

35

40

<210> 61

<211> 53

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(53)

<223> Barstar binding barnase-p3 fusion insert

<400> 61

Leu Gln Ser Ser Gly Gln Asp Ser Gln Arg Glu His Ala Ser His Thr

1

5

10

15

Ala Glu Asp Asp Cys Glu Asp Gln Thr Arg Ile His Gln His Ile Arg

20

25

Glu Val Asp Phe Val Asp Thr Pro Gln Glu Val Asp Asp Cys Arg Ala

35 40 45

Ala Leu Ser Ser Ala

50

<210> 62

<211> 33

<212> PRT

<213> Artificial

<220>

√221> MISC_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 62

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

1 5 10 15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

20 25 30

Ala

<210> 63

<211> 9

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(9)

<223> Barstar binding barnase-p3 fusion insert

<400> 63

Leu Gln Ser Ser Gly Val Arg Pro Ala

1 5

<210> 64

<211> 44

<212> PRT

<213> Artificial

```
<220>
```

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

20

25

30

(112

Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala

35

40

<210> 65

<211> 30

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(30)

<223> Barstar binding barnase-p3 fusion insert

<400> 65

Leu Gln Ser Ser Gly Thr Glu Val Asp Arg Gly Asn Gln Gln His Asp

1

5

10

15

Thr Asn Asp Arg Asp Phe Thr His Thr Pro Leu Ser Ser Ala

20

25

30

LL -210> 6

<211> 36

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 66

Leu Gln Ser Ser Gly Val Ala Gln Gly Ser Ser Ala Ser Val Asp Val

1

5

10

Thr Ala Thr Asn Ala Val Leu Ser Ala Asp Ser Leu Ser Leu Gly Gly

25

20

30

Gly Glu Pro Ala

35

<210> 67

<211> 19

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(19)

<223> Barstar binding barnase-p3 fusion insert

<400> 67

Leu Gln Ser Ser Gly Gly Ala Val Ala Val Thr Pro Gly Pro Val Leu

1 5 10 15

Ser Ser Ala

```
<210> 68
```

<220>

012

<400> 68

Leu Gln Ser Ser Gly His Cys Arg Gly Lys Pro Val Leu Cys Thr His

1

5

10

15

Thr Ala

<210> 69

<211> 9

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(9)

<223> Barstar binding barnase-p3 fusion insert

<400> 69

Leu Gln Ser Ser Gly Val Arg Pro Ala

5

1

<210> 70

<211> 36

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 70

Leu Gln Ser Ser Gly Glu Pro Ala Pro Ala His Glu Ala Lys Pro Thr

1

5

10

Glu Ala Pro Val Ala Lys Ala Glu Ala Lys Pro Glu Thr Pro Ala His

20 25 30

Leu Ser Ser Ala

35

<210> 71

<211> 33

<212> PRT

√213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 71

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

1 5 10 15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

20 25 30

Ala

<210> 72

<211> 36

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 72

Leu Gln Ser Ser Gly Val Val Asp Trp Ala Lys Met Arg Glu Ile Ala

1 5 10 15

Asp Ser Ile Gly Ala Tyr Leu Phe Val Asp Met Ala His Val Ala Ala
20 25 30

```
Leu Ser Ser Ala
```

35

<210> 73

<211> 117

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(117)

<223> Vector pK1 polylinker sequence

012

<400> 73

aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgaggggcc 60

cacgaaggag gtggggtacc cggttccgag ggtggttccg gttccggtga ttttgat 117

<210> 74

<211> 39

<212> PRT

<213> Artificial

<220>

<221> MISC_FEATURE

```
<222> (1)..(39)
```

<223> Polypeptide encoded by pK1 vector polylinker sequence

<400> 74

Asn Ala Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu

1 5 10 15

Gly Arg Gly Ala His Glu Gly Gly Gly Val Pro Gly Ser Glu Gly Gly
20 25 30

012

Ser Gly Ser Gly Asp Phe Asp

35

<210> 75

<211> 117

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(117)

<223> vector pK2 polylinker sequence

```
<400> 75
```

aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgagggccc 60

acgaagcagc tggggtaccg gttccgaggg tggttccggt tccggtgatt ttgatta 117

<210> 76

<211> 39

<212> PRT

<213> Artificial

<220:

<221> MISC_FEATURE

<222> (1)..(39)

<223> Polypeptide sequence encoded by vector pK2 polylinker region.

<220>

<221> MISC_FEATURE

<222> (38)..(38)

<223> X represents a TGA stop codon

<220>

<221> MISC_FEATURE

<222> (36)..(36)

<223> X represents a stop codon (TGA)

Asn Ala Gly Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr/Ile Glu

1

5 .

10

15

Gly Arg Gly Pro Thr Lys Gln Leu Gly Tyr Arg Phe Arg Gly Trp Phe

20

25

30

Arg Phe Arg Xaa Phe Xaa Leu

35

<210> 77

<211> 35

<212> DNA

<213> Artificial

<220>

<221> misc_feature

<222> (1)..(35)

<223> Sequence of the junction region between Barnase and p3 in recombinant fusion vector fd-3.

atcagactgc aggcggtgcg gccgcagaaa ctgtt

35

<210> 78

<211> 11

PRT <212>>>

MISC_FEATURE

<222> (1)..(11)

Amino acid sequence about the junction of barnase and p3 coding r egions of recombinant fusion yector fd-3.

<400> 78

Ile Arg Leu Gln Ala Ala Ala Ala Glu Thr Val

1

5

10

1